

ENTERED

PCT09

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/701,014

DATE: 03/05/2002 TIME: 14:12:21

Input Set : A:\sequence list.txt

Output Set: N:\CRF3\03052002\I701014.raw

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3 <110> APPLICANT: Mitraphanous, Kyrlacos
 4
         Patil, Deva
         Kingsman, Alan
 5
         Kingsman, Susan
         Ellard, Fiona
 9 <120> TITLE OF INVENTION: Retroviral Delivery System
11 <130> FILE REFERENCE: 550-184
13 <140> CURRENT APPLICATION NUMBER: 09/701,014
14 <141> CURRENT FILING DATE: 2001-05-24
16 <150> PRIOR APPLICATION NUMBER: PCT/GB99/01607
17 <151> PRIOR FILING DATE: 1999-05-21
19 <150> PRIOR APPLICATION NUMBER: GB 9811153.7
20 <151> PRIOR FILING DATE: 1998-07-17
22 <160> NUMBER OF SEQ ID NOS: 4
24 <170> SOFTWARE: PatentIn Ver. 2.0
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 1650
28 <212> TYPE: DNA
29 <213> ORGANISM: Artificial Sequence
31 <220> FEATURE:
32 <223> OTHER INFORMATION: Description of Artificial Sequence: rabies virus strain ERA
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37 acatcacete agetgeceaa acaatttggt agtggaggae gaaggatgea ecaacetgte 180
38 agggttctcc tacatggaac ttaaagttgg atacatctta gccataaaaa tgaacgggtt 240
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40 aaccacgttc aaaagaaagc atttccgccc aacaccagat gcatgtagag ccgcgtacaa 360
41 ctggaagatg geeggtgaee eeagatatga agagteteta eacaateegt accetgaeta 420
42 ccgctggctt cgaactgtaa aaaccaccaa ggagtctctc gttatcatat ctccaagtgt 480
43 agcagatttg gacccatatg acagatccct tcactcgagg gtcttcccta gcgggaagtg 540
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45 gcccgagaat ccgagactag ggatgtcttg tgacattttt accaatagta gagggaagag 660
46 agcatccaaa gggagtgaga cttgcggctt tgtagatgaa agaggcctat ataagtcttt 720
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48 atgggtcgcg atgcaaacat caaatgaaac caaatggtgc cctcccgatc agttggtgaa 840
49 cctgcacgac tttcgctcag acgaaattga gcaccttgtt gtagaggagt tggtcaggaa 900
50 gagagaggag tgtctggatg cactagagtc catcatgaca accaagtcag tgagtttcag 960
51 acgtctcagt catttaagaa aacttgtccc tgggtttgga aaagcatata ccatattcaa 1020
52 caagacettg atggaageeg atgeteacta caagteagte agaaettgga atgagateet 1080
 53 cccttcaaaa gggtgtttaa gagttggggg gaggtgtcat cctcatgtga acggggtgtt 1140
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 55 cctcctccag caacatatgg agttgttgga atcctcggtt atcccccttg tgcaccccct 1260
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56 ggcagacccg tctaccgttt tcaaggacgg tgacgaggct gaggattttg ttgaagttca 1320
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 58 gtatgtatta ctgagtgcag gggccctgac tgccttgatg ttgataattt tcctgatgac 1440
 59 atgttgtaga agagtcaatc gatcagaacc tacgcaacac aatctcagag ggacagggag 1500
 60 ggaggtgtca gtcactcccc aaagcgggaa gatcatatct tcatgggaat cacacaagag 1560
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                  20
  79 Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val
                                   40
              35
  80
 82 Val Glu Asp Glu Gly Cys Thr Asn Leu Ser Gly Phe Ser Tyr Met Glu
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                                                   60
  83
85 Leu Lys Val Gly Tyr Ile Leu Ala Ile Lys Met Asn Gly Phe Thr Cys
<del>≐</del> 86
88 Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr
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                      85
91 Val Thr Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Pro Asp Ala
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                 100
  94 Cys Arg Ala Ala Tyr Asn Trp Lys Met Ala Gly Asp Pro Arg Tyr Glu
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             115
  97 Glu Ser Leu His Asn Pro Tyr Pro Asp Tyr Arg Trp Leu Arg Thr Val
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  100 Lys Thr Thr Lys Glu Ser Leu Val Ile Ile Ser Pro Ser Val Ala Asp
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                           150
  101 145
  103 Leu Asp Pro Tyr Asp Arg Ser Leu His Ser Arg Val Phe Pro Ser Gly
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                       165
  106 Lys Cys Ser Gly Val Ala Val Ser Ser Thr Tyr Cys Ser Thr Asn His
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                   180
  109 Asp Tyr Thr Ile Trp Met Pro Glu Asn Pro Arg Leu Gly Met Ser Cys
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               195
  112 Asp Ile Phe Thr Asn Ser Arg Gly Lys Arg Ala Ser Lys Gly Ser Glu
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  115 Thr Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly
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                           230
  116 225
  118 Ala Cys Lys Leu Lys Leu Cys Gly Val Leu Gly Leu Arg Leu Met Asp
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  121 Gly Thr Trp Val Ala Met Gln Thr Ser Asn Glu Thr Lys Trp Cys Pro
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  124 Pro Asp Gln Leu Val Asn Leu His Asp Phe Arg Ser Asp Glu Ile Glu
```

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	13N	Δla	Leu	Glu	Ser	Ile	Met	Thr	Thr	Lys	Ser	Val	Ser	Phe	Arg	Arg	Leu
	121	305					310					315					320
	133	Ser	His	Leu	Arq	Lys	Leu	Val	Pro	Gly	Phe	Gly	Lys	Ala	Tyr	Thr	Ile
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	136	Phe	Asn	Lys	Thr	Leu	Met	Glu	Ala	Asp	Ala	His	Tyr	Lys	Ser	Val	Arg
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	139	Thr	Trp	Asn	Glu	Ile	Leu	Pro	Ser	Lys	Gly	Cys	Leu	Arg	Val	Gly	Gly
	140			355					360					365			
	142	Arq	Cys	His	Pro	His	Val	Asn	Gly	Val	Phe	Phe	Asn	Gly	Ile	Ile	Leu
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	145	Gly	Pro	Asp	Gly	Asn	Val	Leu	Ile	Pro	Glu	Met	Gln	Ser	Ser	Leu	Leu
	116	325					.390					395					400
	148	Gln	Gln	His	Met	Glu	Leu	Leu	Glu	Ser	Ser	Val	Iļe	Pro	Leu	Val	His
	1/10					405					410					410	
	151	Pro	Leu	Ala	Asp	Pro	Ser	Thr	Val	Phe	Lys	Asp	Gly	Asp	Glu	Ala	Glu
	154	Asp	Phe	Val	Glu	Val	His	Leu	Pro	Asp	Val	His	Asn	Gln	Val	Ser	GIY
-	122			471					710								
	157	Val	Asp	Leu	Gly	Leu	Pro	Asn	Trp	Gly	Lys	Tyr	Val	Leu	Leu	Ser	Ala
	158		450					455					460				
	160	Gly	Ala	Leu	Thr	Ala	Leu	Met	Leu	Ile	Ile	Phe	Leu	Met	Thr	Cys	Cys
	161	465					470					475		_	_	a 1	480
	163	Arg	Arg	Val	Asn	Arg	Ser	Glu	Pro	Thr	Gln	His	Asn	Leu	Arg	GTÄ	Thr
	164					485				_	490		_	- 1.	т1.	495	
	166	Gly	Arg	Glu	Val	Ser	Val	Thr	Pro	Gln	Ser	GLY	Lys	тте	ille	ser	Ser
	167				500					505			_		510		
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VERIFICATION SUMMARY

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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date